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## APPL PARTS

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Misc. Incoming Letter

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Notice of Appeal

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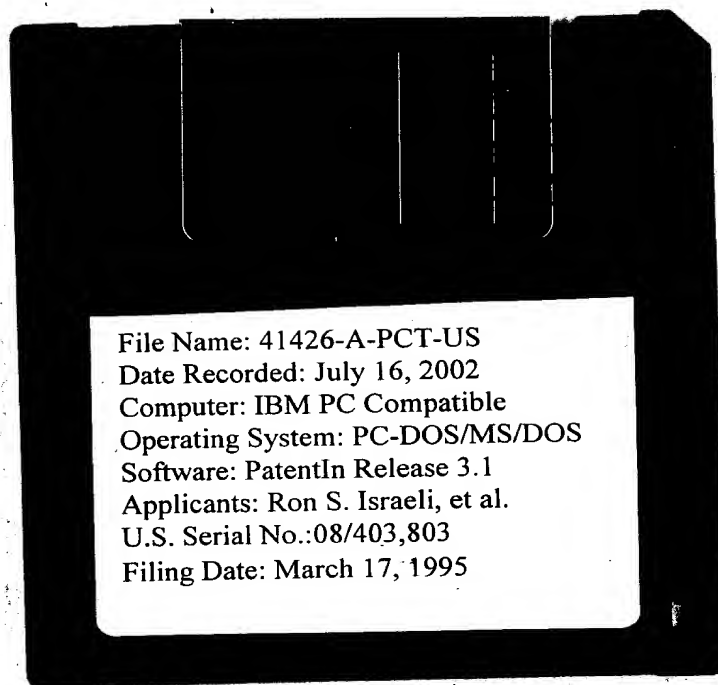
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Team: OIPEBackFileIndexing  
Dossier: 08403803

Legal Date: 08-07-2002

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53 atcaattgct ctgggaaaat tgtaattgcc agatatggga aagttttcag aggaaataag 900
55 gttaaaaatg cccagctggc aggggccaaa ggagtcattc tctactccga cctgctgac 960
57 tactttgctc ctggggtgaa gtcctatcca gatggttgga atcttctg aggtggtgtc 1020
59 cagcgtggaa atatcctaaa tctgaatggt gcaggagacc ctctcacacc aggttaccca 1080
61 gcaaataaat atgcttatag gcgtggaatt gcagaggctg ttggtcttcc aagtattcct 1140
63 gttcatccaa ttggatacta tgatgcacag aagctcctag aaaaaatggg tggctcagca 1200
65 ccaccagata gcagctggag aggaagtctc aaagtgcctt acaatgttg accctggttt 1260
67 actggaaaact tttctacaca aaaagtcaag atgcacatcc actctacca tgaagtgaca 1320
69 agaatttaca atgtgatagg tactctcaga ggagcagtg aaccagacag atatgtcatt 1380
71 ctgggagggtc accgggactc atgggtgttt ggtggtattg accctcagag tggagcagct 1440
73 gttgttcatg aaattgtgag gagctttgga acactgaaaa aggaagggtg gagacctaga 1500
75 agaacaattt tgtttgcaag ctgggatgca gaagaatttg gtcttcttgg ttctactgag 1560
77 tgggcagagg agaattcaag actccttcaa gagcgtggcg tggcttatat taatgctgac 1620
79 tcatctatag aaggaaacta cactctgaga gttgattgta caccgctgat gtacagcttg 1680
81 gtacacaacc taacaaaaga gctgaaaagc cctgatgaag gctttgaagg caaatctctt 1740

```

## RAW SEQUENCE LISTING

DATE: 08/07/2002

PATENT APPLICATION: US/08/403,803C

TIME: 10:24:18

Input Set : A:\1747-41426-A-PCT-US.txt

Output Set: N:\CRF3\08072002\H403803C.raw

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83 tatgaaagtt ggactaaaaa aagtccttcc ccagagttca gtggcatgcc caggataagc 1800
85 aaattgggat ctggaaatga ttttgagggtg ttcttccaac gacttggaat tgcttcaggc 1860
87 agagcacggt atactaaaaa ttgggaaaca aacaaattca gcggctatcc actgtatcac 1920
89 agtgtctatg aaacatatga gttgtgtgaa aagttttatg atccaatgtt taaatatcac 1980
91 ctcaactgtg cccaggttcg aggagggatg gtgtttgagc tagccaattc catagtgtct 2040
93 ccttttgatt gtgcagatta tgctgtagtt ttaagaaagt atgctgacaa aatctacagt 2100
95 atttttatga aacatccaca ggaaatgaag acatacagtg tatcatttga ttcacttttt 2160
97 tctgcagtaa agaattttac agaaattgct tccaagttca gtgagagact ccaggacttt 2220
99 gacaaaagca acccaatagt attaagaatg atgaatgatc aactcatgtt tctggaaaga 2280
101 gcattttattg atccattagg gttaccagac aggccttttt ataggcatgt catctatgct 2340
103 ccaagcagcc acaacaagta tgcaggggag tcattcccag gaatttatga tgctctgttt 2400
105 gatattgaaa gcaaagtgga ccctccaag gcctggggag aagtgaagag acagatttat 2460
107 gttgcagcct tcacagtgca ggcagctgca gagactttga gtgaagtagc ctaagaggat 2520
109 tcttttagaga atcctgtattg aatttgtgtg gtatgtcact cagaaagaat cgtaatgggt 2580
111 atattgataa attttaaaat tggatatatt gaaataaagt tgaatattat atataaaaaa 2640
113 aaaaaaaaaa aaa 2653
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117 <211> LENGTH: 750
118 <212> TYPE: PRT
119 <213> ORGANISM: Human
121 <400> SEQUENCE: 2
123 Met Trp Asn Leu Leu His Glu Thr Asp Ser Ala Val Ala Thr Ala Arg
124 1 5 10 15
127 Arg Pro Arg Trp Leu Cys Ala Gly Ala Leu Val Leu Ala Gly Gly Phe
128 20 25 30
131 Phe Leu Leu Gly Phe Leu Phe Gly Trp Phe Ile Lys Ser Ser Asn Glu
132 35 40 45
135 Ala Thr Asn Ile Thr Pro Lys His Asn Met Lys Ala Phe Leu Asp Glu
136 50 55 60
139 Leu Lys Ala Glu Asn Ile Lys Lys Phe Leu Tyr Asn Phe Thr Gln Ile
140 65 70 75 80
143 Pro His Leu Ala Gly Thr Glu Gln Asn Phe Gln Leu Ala Lys Gln Ile
144 85 90 95
147 Gln Ser Gln Trp Lys Glu Phe Gly Leu Asp Ser Val Glu Leu Ala His
148 100 105 110
151 Tyr Asp Val Leu Leu Ser Tyr Pro Asn Lys Thr His Pro Asn Tyr Ile
152 115 120 125
155 Ser Ile Ile Asn Glu Asp Gly Asn Glu Ile Phe Asn Thr Ser Leu Phe
156 130 135 140
159 Glu Pro Pro Pro Pro Gly Tyr Glu Asn Val Ser Asp Ile Val Pro Pro
160 145 150 155 160
163 Phe Ser Ala Phe Ser Pro Gln Gly Met Pro Glu Gly Asp Leu Val Tyr
164 165 170 175
167 Val Asn Tyr Ala Arg Thr Glu Asp Phe Phe Lys Leu Glu Arg Asp Met
168 180 185 190
171 Lys Ile Asn Cys Ser Gly Lys Ile Val Ile Ala Arg Tyr Gly Lys Val
172 195 200 205
175 Phe Arg Gly Asn Lys Val Lys Asn Ala Gln Leu Ala Gly Ala Lys Gly
176 210 215 220

```

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179 Val Ile Leu Tyr Ser Asp Pro Ala Asp Tyr Phe Ala Pro Gly Val Lys
180 225 230 235 240
183 Ser Tyr Pro Asp Gly Trp Asn Leu Pro Gly Gly Val Gln Arg Gly
184 245 250 255
187 Asn Ile Leu Asn Leu Asn Gly Ala Gly Asp Pro Leu Thr Pro Gly Tyr
188 260 265 270
191 Pro Ala Asn Glu Tyr Ala Tyr Arg Gly Ile Ala Glu Ala Val Gly
192 275 280 285
195 Leu Pro Ser Ile Pro Val His Pro Ile Gly Tyr Tyr Asp Ala Gln Lys
196 290 295 300
199 Leu Leu Glu Lys Met Gly Gly Ser Ala Pro Pro Asp Ser Ser Trp Arg
200 305 310 315 320
203 Gly Ser Leu Lys Val Pro Tyr Asn Val Gly Pro Gly Phe Thr Gly Asn
204 325 330 335
207 Phe Ser Thr Gln Lys Val Lys Met His Ile His Ser Thr Asn Glu Val
208 340 345 350
211 Thr Arg Ile Tyr Asn Val Ile Gly Thr Leu Arg Gly Ala Val Glu Pro
212 355 360 365
215 Asp Arg Tyr Val Ile Leu Gly Gly His Arg Asp Ser Trp Val Phe Gly
216 370 375 380
219 Gly Ile Asp Pro Gln Ser Gly Ala Ala Val Val His Glu Ile Val Arg
220 385 390 395 400
223 Ser Phe Gly Thr Leu Lys Lys Glu Gly Trp Arg Pro Arg Arg Thr Ile
224 405 410 415
227 Leu Phe Ala Ser Trp Asp Ala Glu Glu Phe Gly Leu Leu Gly Ser Thr
228 420 425 430
231 Glu Trp Ala Glu Glu Asn Ser Arg Leu Leu Gln Glu Arg Gly Val Ala
232 435 440 445
235 Tyr Ile Asn Ala Asp Ser Ser Ile Glu Gly Asn Tyr Thr Leu Arg Val
236 450 455 460
239 Asp Cys Thr Pro Leu Met Tyr Ser Leu Val His Asn Leu Thr Lys Glu
240 465 470 475 480
243 Leu Lys Ser Pro Asp Glu Gly Phe Glu Gly Lys Ser Leu Tyr Glu Ser
244 485 490 495
247 Trp Thr Lys Lys Ser Pro Ser Pro Glu Phe Ser Gly Met Pro Arg Ile
248 500 505 510
251 Ser Lys Leu Gly Ser Gly Asn Asp Phe Glu Val Phe Phe Gln Arg Leu
252 515 520 525
255 Lys Ile Ala Ser Gly Arg Ala Arg Tyr Thr Lys Asn Trp Glu Thr Asn
256 530 535 540
259 Lys Phe Ser Gly Tyr Pro Leu Tyr His Ser Val Tyr Glu Thr Tyr Glu
260 545 550 555 560
263 Leu Val Glu Lys Phe Tyr Asp Pro Met Phe Lys Tyr His Leu Thr Val
264 565 570 575
267 Ala Gln Val Arg Gly Gly Met Val Phe Glu Leu Ala Asn Ser Ile Val
268 580 585 590
271 Leu Pro Phe Asp Cys Arg Asp Tyr Ala Val Val Leu Arg Lys Tyr Ala
272 595 600 605
275 Asp Lys Ile Tyr Ser Ile Ser Met Lys His Pro Gln Glu Met Lys Thr

```

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Input Set : A:\1747-41426-A-PCT-US.txt

Output Set: N:\CRF3\08072002\H403803C.raw

```

276      610      615      620
279 Tyr Ser Val Ser Phe Asp Ser Leu Phe Ser Ala Val Lys Asn Phe Thr
280 625      630      635      640
283 Glu Ile Ala Ser Lys Phe Ser Glu Arg Leu Gln Asp Phe Asp Lys Ser
284      645      650      655
287 Asn Pro Ile Val Leu Arg Met Met Asn Asp Gln Leu Met Phe Leu Glu
288      660      665      670
291 Arg Ala Phe Ile Asp Pro Leu Gly Leu Pro Asp Arg Pro Phe Tyr Arg
292      675      680      685
295 His Val Ile Tyr Ala Pro Ser Ser His Asn Lys Tyr Ala Gly Glu Ser
296      690      695      700
299 Phe Pro Gly Ile Tyr Asp Ala Leu Phe Asp Ile Glu Ser Lys Val Asp
300 705      710      715      720
303 Pro Ser Lys Ala Trp Gly Glu Val Lys Arg Gln Ile Tyr Val Ala Ala
304      725      730      735
307 Phe Thr Val Gln Ala Ala Ala Glu Thr Leu Ser Glu Val Ala
308      740      745      750
311 <210> SEQ ID NO: 3
312 <211> LENGTH: 8
313 <212> TYPE: PRT
314 <213> ORGANISM: Human
316 <400> SEQUENCE: 3
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319 1      5
322 <210> SEQ ID NO: 4
323 <211> LENGTH: 15
324 <212> TYPE: PRT
325 <213> ORGANISM: Human
327 <220> FEATURE:
328 <221> NAME/KEY: MISC_FEATURE
329 <222> LOCATION: (6)..(7)
330 <223> OTHER INFORMATION: Xaa=unknown
333 <400> SEQUENCE: 4
W--> 335 Ser Tyr Pro Asp Gly Xaa Xaa Leu Pro Gly Gly Gly Val Gln Arg
336 1      5      10      15
339 <210> SEQ ID NO: 5
340 <211> LENGTH: 7
341 <212> TYPE: PRT
342 <213> ORGANISM: Human
344 <400> SEQUENCE: 5
346 Phe Tyr Asp Pro Met Phe Lys
347 1      5
350 <210> SEQ ID NO: 6
351 <211> LENGTH: 9
352 <212> TYPE: PRT
353 <213> ORGANISM: Human
355 <400> SEQUENCE: 6
357 Ile Tyr Asn Val Ile Gly Thr Leu Lys
358 1      5

```